STIC-Biotech/Ch mLib

Fr m: Sent: Subject:

STIC-ILL Wednesday, March 06, 2002 9:19 AM STIC-Biotech/ChemLib

FW: sequence search request

----Original Message-

From:

Chernyshev, Olga

Sent:

Wednesday, March 06, 2002 9:17 AM

To:

STIC-ILL

Subject:

sequence search request

US case 09/853,753 Please search SEQ ID NO:1 and 2. Thank you very much!

Olga N. Chernyshev AŬ1646 CM1 8D06 *305-1003* mail box 10C01

> **Edward Hart** Technicai Info. Specialist STIC/Biotech CMI 6B02 Tel: 305-9203

Searcher:
Phone:
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TYPE OF SEARCH:
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Database

GenEmbl:*

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Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

stal number of hits satisfying chosen parameters:

2944280

1472140 seqs, 8248589755 residues

IDENTITY_NUC
Gapop 10.0 , Gapext 1.0

Scoring table: Sequence: Perfect score:

US-09-853-753-1 2297 1-ggctgagggagtggagggg.....aaggttaaaccggtttctct 2297

Run on:

OM nucleic - nucleic search, using sw model

GenCore version 4.5 Copyright (c) 1993 - 2000 Com

Compugen Ltd

March 8,

2002, 08:51:10 ; Search time 2236.97 Seconds (without alignments)
16939.888 Million cell updates/sec

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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Bech-Hansen, N.T., Maybaum, T.A. and Naylor, M.J.

Birect Submission

Submitted (10-APR-2000) Medical Genetics, University of Calgary,

3330 Hospital Dr. N.W., Calgary, AB T2N 4N1, Canada
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1560 1560	ccagcgtggcccagcacgtggtgtttggcctgcagatggactgacctggccagagggggg	1501 1501	Db Qy
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1140 1140	tgggccgcctcttcctcttccgcaacccgtggtgctgcgactgccgtctggagtggctga 	1081	ga ED
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600	TCAGCGTGCCCGAGCGCCTCCTGGCCGAACTGCCGGCCCTGCGCGAACTCGCCGTTCG	. 541	3 .

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Nat. Gene
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AJ278865
AJ278865.1 G:
CLRP gene; lev
                                                                                                                                       Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 2713)

Pusch,C.M., Zeitz,C., Brandau,O., Pesch,K., Achatz,H., Feil,S., Scharfe,C., Maurer,J., Jacobi,F.K., Pinckers,A., Andreasson,S., Hardcastle,A., Wissinger,B., Berger,W. and Meindl,A.

The complete form of X-linked congenital stationary night blinds is caused by mutations in a gene encoding a leucine-rich repeat
                                                                                                                                                                                                                                           human.
                                                    Direct Submission
Submitted (23-AUG-2000) Medical Genetics, Ludwigs Maximilians
University, Goethestr. 29, Munich D-80336, GERMANY
                                                                                                                                                                                                                      Homo sapiens
Eukaryota; Metazoa;
                                                                                      Pusch, C.
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Primates;
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NLTTSSPGPSPBPAATTVSRFESSLEXKLLAPRVPEEAANTTGGLANASLSDSLSSRG
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457. .1902
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100.0%; F
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                                            Score 2241; DB 9;
Pred. No. 3.3e-252;
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